CLAIMS

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- 1. A process of obtaining a species which inhibits the activity of a polypeptide, comprising:
 - a. contacting an active polypeptide fragment of the polypeptide with the species;
 - determining if the species interacts with the active polypeptide fragment; and
 - c. determining if the species inhibits the activity of the polypeptide.
- 2. The process of claim 1 wherein the species is a chemical compound.
- 3. The process of claim 1 wherein the species is a polypeptide.
- 4. The process of claim 1 wherein the species is a polynucleotide.
- 5. The process of claim 1 wherein the species is imprinted on a microarray.
- 6. The process of claim 1 wherein the polypeptide is known to have an effect in a disease state or condition.
 - 7. The process of claim 1 wherein the species is a chemical compound imprinted on a microarray and the active polypeptide fragment is a dominant polypeptide fragment.
 - 8. The process of claim 1 wherein the species is a chemical compound imprinted on a microarray and the active polypeptide fragment is a polypeptide fragment comprising an active site of one or more polypeptide of interest.

- 9. A process of obtaining a species that modulates a phenotype comprising:
 - (a) contacting a microarray comprising an array of species with a plurality of active polypeptide fragments;
 - (b) examining the microarray for a positively reacting species;
 - (c) obtaining one or more species which react positively; and
 - (d) examining the effect of the species of step (c) on a cell exhibiting the phenotype.
- 10. The process of claim 9 further comprising:
 - (e) identifying the active polypeptide fragment which caused the species to react;
 - (f) obtaining a parent polypeptide; and
 - (g) examining the effect of the species on the parent polypeptide.
- 15 11. The process of claim 9 further comprising:
 - (e) identifying the active polypeptide fragment which caused the species to react;
 - (f) obtaining a parent polypeptide; and
 - (g) examining the effect of the parent polypeptide on the phenotype.
 - 12. The process of claim 9 wherein one or more of the active polypeptide fragments of step (a) are associated with the phenotype exhibited by the cell of step (d).
- 13. The process of claim 9 wherein the active polypeptide fragments of step (a) are tagged.
 - 14. The process of claim 9 wherein the species imprinted on the microarray are chemical compounds.

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- 15. The process of claim 9 wherein the species imprinted on the microarray are polypeptides.
- 16. The process of claim 9 wherein the species imprinted on the microarray are polynucleotides.
- 17. A kit for performing the process of claim 9 which comprises:
 - a. a microarray; and
 - b. a plurality of active polypeptide fragments.

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- 18. The kit of claim 17 wherein the plurality of active polypeptide fragments are associated with a phenotype.
- 19. A process of obtaining a small molecule polypeptide inhibitor comprising:
 - a) screening a plurality of small molecules with a polypeptide probe;
 - b) screening the plurality of small molecules of step a) with a second polypeptide probe comprising the polypeptide probe of step a) further comprising one or more amino acid mutation;
 - c) comparing the small molecule binding profile of the probe of step b) to that of the probe of step a); and
 - d) identifying one or more small molecules from the plurality of small molecules that bind to the probe of step a) but not to the probe of step b).
- 20. The process of claim 19 wherein the plurality of small molecules is imprinted on a microarray.
- 21. The process of claim 19 wherein the mutation of step b) is randomly generated.
- 22. The process of claim 19 wherein the mutation of step b) is in a region of the polypeptide known to be involved in a biological activity.

23. The process of claim 22 wherein the biological activity is selected from the biological activity group consisting of: enzymatic activity, protein-protein interaction, protein-DNA interaction, protein-chemical interaction, protein-carbohydrate interaction, protein modification, localization signal, ATP/GTP carrying site, and ion binding.

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24. The process of claim 22 wherein the mutation of step b) is in a region of the polypeptide known to be involved in an enzymatic activity.